FIGURE 1

CAGAGAAAACTCCTCCAAAAGCAGCTCTCACTATCAGAAAACCCAACTAC AGTTGTGAACGCCTTCATTTTCTGCCTGAGGTCTCAGTCCGTCGGCCCAG ACTGAAGTGCAGTGGCACAATCATAGCTCGCTGCAGCCTCGACCTTCCAG GCTTAAACGATTCTCCCACCTCAGCCTCTCGAGTTGCTGGGACCACAGGT CACTGTGCATGGCATCATCCTGGCCCCCTCTAGAGCTCCAATCCTCCAAC CAGAGCCAGCTCTTCCCTCAAAATGCTACGGCCTGTGACAATGCTCCAGA AGCCTGGGACCTGCTGCACAGAGTGCTGCCGACATTTATCATCTCCATCT GTTTCTTCGGCCTCCTAGGGAACCTTTTTGTCCTGTTGGTCTTCCTCCTG CCCCGGCGCAACTGAACGTGGCAGAAATCTACCTGGCCAACCTGGCAGC CTCTGATCTGGTGTTTGTCTTGGGCTTGCCCTTCTGGGCAGAGAATATCT GGAACCAGTTTAACTGGCCTTTCGGAGCCCTCCTCTGCCGTGTCATCAAC GGGGTCATCAAGGCCAATTTGTTCATCAGCATCTTCCTGGTGGTGGCCAT CAGCCAGGACCGCTACCGCGTGCTGGTGCACCCTATGGCCAGCGGAAGGC AGCAGCGGCGGAGGCCAGGCCCGGGTCACCTGCGTGCTCATCTGGGTTGTG CCCAGATCTGAACATCACCGCCTGCATCCTGCTCCTCCCCCATGAGGCCT GGCACTTTGCAAGGATTGTGGAGTTAAATATTCTGGGTTTCCTCCTACCA CTGGCTGCGATCGTCTTCTTCAACTACCACATCCTGGCCTCCCTGCGAAC GCGGGAGGAGGTCAGCAGGACAAGGTGCGGGGGCCGCAAGGATAGCAAGA CCACAGCGCTGATCCTCACGCTCGTGGTTGCCTTCCTGGTCTGCTGGGCC CCTTACCACTTCTTTGCCTTCCTGGAATTCTTATTCCAGGTGCAAGCAGT CCGAGGCTGCTTTTGGGAGGACTTCATTGACCTGGGCCTGCAATTGGCCA ACTTCTTTGCCTTCACTAACAGCTCCCTGAATCCAGTAATTTATGTCTTT GTGGGCCGGCTCTTCAGGACCAAGGTCTGGGAACTTTATAAACAATGCAC AAAAAA SEO.ID.NO.:1.

FIGURE 2

MET Ala Ser Ser Trp Pro Pro Leu Glu Leu Gln Ser Ser Asn Gln Ser Gln Leu Phe Pro Gln Asn Ala Thr Ala Cys Asp Asn Ala Pro Glu Ala Trp Asp Leu Leu His Arg Val Leu Pro Thr Phe Ile Ile Ser Ile Cys Phe Phe Gly Leu Leu Gly Asn Leu Phe Val Leu Leu Val Phe Leu Leu Pro Arg Arg Gln Leu Asn Val Ala Glu Ile Tyr Leu Ala Asn Leu Ala Ala Ser Asp Leu Val Phe Val Leu Gly Leu Pro Phe Trp Ala Glu Asn Ile Trp Asn Gln Phe Asn Trp Pro Phe Gly Ala Leu Leu Cys Arg Val Ile Asn Gly Val Ile Lys Ala Asn Leu Phe Ile Ser Ile Phe Leu Val Val Ala Ile Ser Gln Asp Arg Tyr Arg Val Leu Val His Pro MET Ala Ser Gly Arg Gln Gln Arg Arg Arg Gln Ala Arg Val Thr Cys Val Leu Ile Trp Val Val Gly Gly Leu Leu Ser Ile Pro Thr Phe Leu Leu Arg Ser Ile Gln Ala Val Pro Asp Leu Asn Ile Thr Ala Cys Ile Leu Leu Pro His Glu Ala Trp His Phe Ala Arg Ile Val Glu Leu Asn Ile Leu Gly Phe Leu Leu Pro Leu Ala Ala Ile Val Phe Phe Asn Tyr His Ile Leu Ala Ser Leu Arg Thr Arg Glu Glu Val Ser Arg Thr Arg Cys Gly Gly Arg Lys Asp Ser Lys Thr Thr Ala Leu Ile Leu Thr Leu Val Val Ala Phe Leu Val Cys Trp Ala Pro Tyr His Phe Phe Ala Phe Leu Glu Phe Leu Phe Gln Val Gln Ala Val Arg Gly Cys Phe Trp Glu Asp Phe Ile Asp Leu Gly Leu Gln Leu Ala Asn Phe Phe Ala Phe Thr Asn Ser Ser Leu Asn Pro Val Ile Tyr Val Phe Val Gly Arg Leu Phe Arg Thr Lys Val Trp Glu Leu Tyr Lys Gln Cys Thr Pro Lys Ser Leu Ala Pro Ile Ser Ser His Arg Lys Glu Ile Phe Gln Leu Phe Trp Arg Asn SEQ.ID.NO.:2.

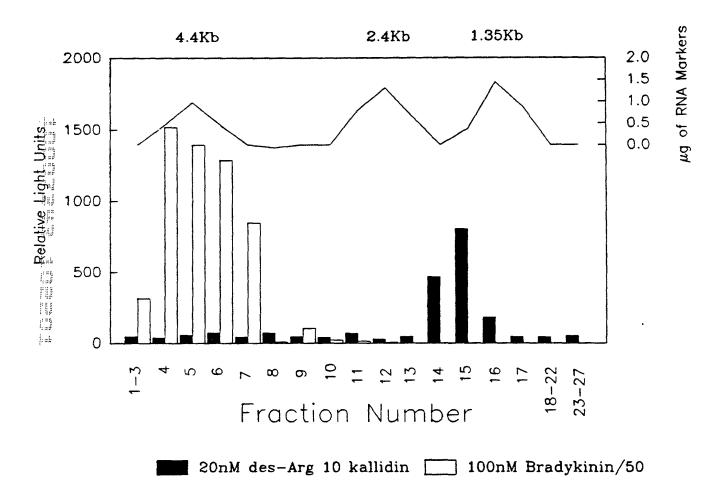
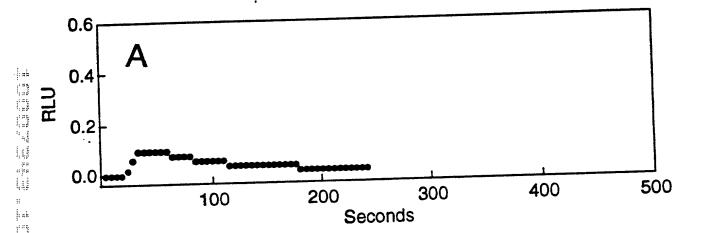


FIGURE 3



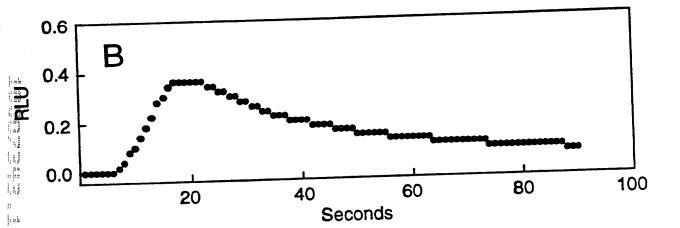


FIGURE 4B

